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RAW SEQUENCE LISTING
 PATENT APPLICATION: US/10/049,404

DATE: 07/11/2002
 TIME: 14:01:57

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3 <110> APPLICANT: Arndt, Michaela
 4 Little, Melvyn
 5 Kypriyanov, Sergey
 6 Krauss, Jurgen
 7 Pfreundschuh, Michael
 9 <120> TITLE OF INVENTION: Fv Antibody Construct Comprising Binding Sites For a CD16
 Receptor and a
 10 CD30 Surface Protein
 12 <130> FILE REFERENCE: 4121-135
 14 <140> CURRENT APPLICATION NUMBER: US 10/049,404
 15 <141> CURRENT FILING DATE: 2002-02-05
 17 <150> PRIOR APPLICATION NUMBER: PCT/DE00/02589
 18 <151> PRIOR FILING DATE: 2000-08-02
 20 <150> PRIOR APPLICATION NUMBER: DE 199 37 264
 21 <151> PRIOR FILING DATE: 1999-08-06
 23 <160> NUMBER OF SEQ ID NOS: 11
 25 <170> SOFTWARE: PatentIn version 3.1
 27 <210> SEQ ID NO: 1
 28 <211> LENGTH: 4570
 29 <212> TYPE: DNA
 30 <213> ORGANISM: Artificial Sequence
 32 <220> FEATURE:
 33 <223> OTHER INFORMATION: Synthetic Construct
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 40 atttcacaca gaattcatta aagaggagaa attaaccatg aaatacctat tgcctacggc 180
 42 agccgctggc ttgctgctgc tggcagctca gccgccatgg cgcaggtgca gctgcagcag 240
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 60 gggcaatctc cttaaagttc gatttactcg gcattctacc gatacagtg agtccctgat 780
 62 cgcttcacag gcagtgatc tggaaacagat ttcaacttca ccatcagcaa tgtgcagtct 840
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 66 ggcaccaagc tggaaatcaa acgggctgat gctgcggccg ctggatccga acaaaagctg 960
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96	tgggtgttcg	gtggaggaac	caaactgact	gtcctaggcc	agcccaagtc	tgcggccgct	1860
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104	aatagcgaga	ggcccgaccc	gatcgccctt	cccaacagtt	gcgcagcctg	aatggcgaaat	2100
106	gggacgcgcc	ctgtagcgcc	gcattaagcg	cggcggtgtg	ggtggttacg	cgacgcgtga	2160
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174 agaaagggcgg acagggtatcc ggtaagcggc aggggtcggaa caggagagcg caccagggag 4200
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178 gagcgtcgat ttttgtgatg ctcgtcaggg gggcggagcc tatggaaaaa cgccagcaac 4320
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205 Gly Thr Ser Val Lys Ile Ser Cys Lys Ala Ser Gly Tyr Thr Phe Thr
206 20 25 30
209 Asn Tyr Trp Leu Gly Trp Val Lys Gln Arg Pro Gly His Gly Leu Glu
210 35 40 45
213 Trp Ile Gly Asp Ile Tyr Pro Gly Gly Gly Tyr Thr Asn Tyr Asn Glu
214 50 55 60
217 Lys Phe Lys Gly Lys Ala Thr Val Thr Ala Asp Thr Ser Ser Arg Thr
218 65 70 75 80
221 Ala Tyr Val Gln Val Arg Ser Leu Thr Ser Glu Asp Ser Ala Val Tyr
222 85 90 95
225 Phe Cys Ala Arg Ser Ala Ser Trp Tyr Phe Asp Val Trp Gly Ala Arg
226 100 105 110
229 Thr Thr Val Thr Val Ser Ser Ala Lys Thr Thr Pro Lys Leu Gly Gly
230 115 120 125
233 Asp Ile Glu Leu Thr Gln Ser Pro Lys Phe Met Ser Thr Ser Val Gly
234 130 135 140
237 Asp Arg Val Asn Val Thr Tyr Lys Ala Ser Gln Asn Val Gly Thr Asn
238 145 150 155 160
241 Val Ala Trp Phe Gln Lys Pro Gly Gln Ser Pro Lys Val Leu Ile
242 165 170 175
245 Tyr Ser Ala Ser Tyr Arg Tyr Ser Gly Val Pro Asp Arg Phe Thr Gly
246 180 185 190
249 Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Asn Val Gln Ser
250 195 200 205
253 Glu Asp Leu Ala Glu Tyr Phe Cys Gln Gln Tyr His Thr Tyr Pro Leu
254 210 215 220
257 Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys Arg Ala Asp Ala Ala
258 225 230 235 240
261 Ala Ala Gly Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Asn Ser
262 245 250 255
265 His His His His His His
266 260

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271 <212> TYPE: PRT
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283 Gly Ala Ser Val Lys Met Ser Cys Lys Ala Ser Gly Tyr Thr Phe Thr
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287 Thr Tyr Thr Ile His Trp Val Arg Gln Arg Pro Gly His Asp Leu Glu
288 35 40 45
291 Trp Ile Gly Tyr Ile Asn Pro Ser Ser Gly Tyr Ser Asp Tyr Asn Gln
292 50 55 60
295 Asn Phe Lys Gly Lys Thr Thr Leu Thr Ala Asp Lys Ser Ser Asn Thr
296 65 70 75 80
299 Ala Tyr Met Gln Leu Asn Ser Leu Thr Ser Glu Asp Ser Ala Val Tyr
300 85 90 95
303 Tyr Cys Ala Arg Arg Ala Asp Tyr Gly Asn Tyr Glu Tyr Thr Trp Phe
304 100 105 110
307 Ala Tyr Trp Gly Gln Gly Thr Thr Val Thr Val Ser Ser Ala Lys Thr
308 115 120 125
311 Thr Pro Lys Leu Gly Gly Asp Ile Gln Ala Val Val Thr Gln Glu Ser
312 130 135 140
315 Ala Leu Thr Thr Ser Pro Gly Glu Thr Val Thr Leu Thr Cys Arg Ser
316 145 150 155 160
319 Asn Thr Gly Thr Val Thr Thr Ser Asn Tyr Ala Asn Trp Val Gln Glu
320 165 170 175
323 Lys Pro Asp His Leu Phe Thr Gly Leu Ile Gly His Thr Asn Asn Arg
324 180 185 190
327 Ala Pro Gly Val Pro Ala Arg Phe Ser Gly Ser Leu Ile Gly Asp Lys
328 195 200 205
331 Ala Ala Leu Thr Ile Thr Gly Ala Gln Thr Glu Asp Glu Ala Ile Tyr
332 210 215 220
335 Phe Cys Ala Leu Trp Tyr Asn Asn His Trp Val Phe Gly Gly Gly Thr
336 225 230 235 240
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413 <212> TYPE: DNA
414 <213> ORGANISM: Artificial Sequence
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425 <212> TYPE: DNA
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RAW SEQUENCE LISTING ERROR SUMMARY
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VERIFICATION SUMMARY

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